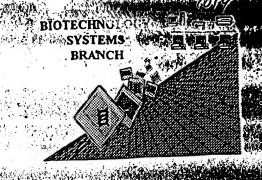
RAW SEQUENCE LISTING



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/966958
Source:	OIPE
Date Processed by STIC:	10/18/01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4 PATENTIN 2.1 c-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 c-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2Kcompliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING DATE: 10/18/2001 PATENT APPLICATION: US/09/966,955 TIME: 09:04:19

Input Set : A:\3053usll.app

Output Set: N:\CRF3\10182001\1966955.raw

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3 <110> APPLICANT: Perez-Villar, Juan J.
              Chang, Han
                                                                       Does Not Comply
              Yang, Wen-Pin
                                                                   Corrected Diskette Needed
             Wu, Yuli
      7
             Whitney, Gena S.
              Kanner, Steven B.
     10 <120> TITLE OF INVENTION: Identification and Cloning of a Full-length Human
     11
              Clnk-related Gene, MIST (Mast Cell Immunoreceptor
     12
              Signal Transducer)
                                                        Dure correct aurent application
     14 <130> FILE REFERENCE: 3053-4113US1
C--> 16 <140> CURRENT APPLICATION NUMBER: US/09/966,955
     17 <141> CURRENT FILING DATE: 2001-09-28
     19 <150> PRIOR APPLICATION NUMBER: 60/237030
     20 <151> PRIOR FILING DATE: 2000-09-29
     22 <160> NUMBER OF SEQ ID NOS: 52
     24 <170> SOFTWARE: PatentIn Ver. 2.1
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ERRORED SEQUENCES

462 <210> SEQ ID NO: 8
463 <211> LENGTH: 21
464 <212> TYPE: DNA
465 <213> ORGANISM: HUMAN
467 <220> FEATURE:
468 <223> OTHER INFORMATION: PRIMER PY474
470 <400> SEQUENCE: 8
E--> 471 tggtacattg gagaatacag

Errored: Input: 21 nucleo tides on field 211 Sequence
Found: 20 nucleo tides Not Comply
Corrected Diskette Needed

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/966,955

DATE: 10/18/2001

TIME: 09:04:20

Input Set : A:\3053usll.app

Output Set: N:\CRF3\10182001\1966955.raw

L:16 M:270 C: Current Application Number differs, Wrong Format

L:471 M:252 E: No. of Seq. differs, <211>LENGTH:Input:21 Found:20 SEQ:8